Moore, et al. (9692-000030) Seq.Listing.ST25 SEQUENCE LISTING <110> Moore, Helen M. Bonazzi, Vivien R. Foster, Leslie L. Gorokhov, Mark Jojic, Olivier Lu, Fu Mason, Douglas Wang, Zhen Y. Wise, Richard E. Zheng, Xiangqun H. Zhong, Fei <120> METHODS FOR IDENTIFYING, VIEWING, AND ANALYZING SYNTENIC AND ORTHOLOGOUS GENOMIC REGIONS BETWEEN TWO OR MORE SPECIES <130> 9692-000030 <160> 21 <170> PatentIn version 3.2 <210> 1 59 <211> <212> DNA <213> Homo sapiens <400> 1 ccagcttcgc ctcaccccat tacagtctca ccacgccccg tcccctctcc gttgagccc 59 <210> 2 58 <211> <212> DNA <213> Homo sapiens <400> 2 cagcttcgcc tcaccccatt acagtctcac cacgccccgt cccctctccg ttgagccc 58 <210> 3 58 <211> <212> DNA <213> Homo sapiens <400> 3 cgcgccttcg cccgggtggg gcgctgcgct gtcagcggcc ttgctgtgtg aggcagaa 58 <210> 4 <211> 58 <212> DNA <213> Homo sapiens <400> 4 cctgcggggg caggggcggg ctggttccct ggccagccat tggcagagtc cgcaggct 58 <210> 5 58 <211> <212> DNA <213> Homo sapiens

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